SEQUENCE LISTING

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<110> Michael F. Thomashow
      Sarah George Fowler
      Jonathan Vogel
      Daniel Zarka
<120> Transcription Factors to Improve Plant Stress Tolerance
<130> 21835-00004
<150> 60/400,777
<151> August 2, 2002
<160> 2
<210> 1
<211> 1308
<212> DNA
<213> Arabidopsis thaliana
<220>
<221>
<222>
<223> This is a cDNA sequence derived from mRNA encoding the putative AP2 domain
transcription factor RAV1 (locus tag: Atlg13260). The RAV1 coding sequence
consists of nucleotides 92 through 1126 as numbered below.
<400> 1
gtatacatat acacaacata attcacaaca caacacaaac acatttctqt tttctccatt
                                                                   60
gtttcaaacc ataaaaaaa acacagatta aatggaatcg agtagcgttg atgagagtac
                                                                   120
tacaagtaca ggttccatct gtgaaacccc ggcgataact ccggcgaaaa agtcgtcggt
                                                                   180
aggtaactta tacaggatgg gaagcggatc aagcgttgtg ttagattcag agaacggcgt
                                                                   240
agaagctgaa tctaggaagc ttccgtcgtc aaaatacaaa ggtgtggtgc cacaaccaaa
                                                                   300
cggaagatgg ggagctcaga tttacgagaa acaccagcgc gtgtggctcg ggacattcaa
                                                                   360
cgaagaagac gaagccgctc gtgcctacga cgtcgcggtt cacaggttcc gtcgccgtga
                                                                   420
                                                                   480
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cgccgtcaca aatttcaaag acgtgaagat ggacgaagac gaggtcgatt tcttgaattc
tcattcgaaa tctgagatcg ttgatatgtt gaggaaacat acttataacg aagagttaga
                                                                       540
gcagagtaaa cggcgtcgta atggtaacgg aaacatgact aggacgttgt taacgtcggg
                                                                       600
gttgagtaat gatggtgttt ctacgacggg gtttagatcg gcggaggcac tgtttgagaa
                                                                       660
agcggtaacg ccaagcgacq ttgggaagct aaaccgtttg gttataccga aacatcacgc
                                                                       720
agagaaacat tttccqttac cqtcaaqtaa cqtttccqtq aaaqqaqtqt tqttqaactt
                                                                       780
tgaggacgtt aacgggaaag tgtggaggtt ccgttactcg tattggaaca gtagtcagag
                                                                       840
ttatgttttg actaaaggtt ggagcaggtt cgttaaggag aagaatctac gtgctggtga
                                                                       900
cgtggttagt ttcagtagat ctaacggtca ggatcaacag ttgtacattg ggtggaagtc
                                                                       960
gagatccggg tcagatttag atgcgggtcg ggttttgaga ttgttcggag ttaacatttc
                                                                       1020
accggagagt tcaagaaacg acgtcgtagg aaacaaaaga gtgaacgata ctgagatgtt
                                                                       1080
atcqttqqtg tgtaqcaaqa agcaacqcat ctttcacqcc tcqtaacaac tcttcttctt
                                                                       1140
tttttttttt tttgttgttt taataatttt taaaaactcc attttcgttt tctttatttg
                                                                       1200
catcggtttc tttcttcttg tttaccaaag gttcatgagt tgtttttgtt gtattgatga
                                                                       1260
actgtaaatt ttatttatag gataaatttt aaaaagggtt acttagat
                                                                       1308
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<210> 2
<211> 1308
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<212> DNA

<213> Arabidopsis thaliana

<220>

<221>

<222>

<223> This is a cDNA sequence derived from mRNA encoding the putative zinc-finger protein Zat12 (locus tag: At5g59820). The Zat12 coding sequence consists of nucleotides 102 through 590 as numbered below.

<400> 2

atcatcacaa cta	actatcac acca	aactca aaaaa	acacaa accac	aagag gatca	tttca 60	
ttttttattg tt	tcgtttta atca	tcatca tcaga	aagaaa aatgg	ttgcg atatc	ggaga 120	0
tcaagtcgac gg	tggatgtc acgg	cggcga attg	ttgat gcttt	tatct agagt	tggac 180	0
aagaaaacgt tg	acggtggc gatc	aaaaac gcgti	ttcac atgta	aaacg tgttt	gaagc 240	0
agtttcattc gt	tccaagcc ttag	gaggtc accgi	gcgag tcaca	agaag cctaa	caacg 300	0
acgctttgtc gto	ctggattg atga	agaagg tgaaa	aacgtc gtcgc	atcct tgtcc	catat 360	0
gtggagtgga gt	ttccgatg ggac	aagctt tggg	aggaca catga	iggaga cacag	gaacg 420	0
agagtggggc tg	ctggtggc gcgt	tggtta cacg	cgcttt gttgc	cggag cccac	ggtga 480	0
ctacgttgaa gaa	aatctagc agtg	ggaaga gagt	ggcttg tttgg	atctg agtct	aggga 540	0
tggtggacaa tt	tgaatctc aagt	tggagc ttgg	aagaac agttt	attga tttta	tttat 600	0
tttccttaaa tt	ttctgaat atat	ttgttt ctct	cattct ttgaa	ttttt cttaa	tattc 660	0
tagattatac ata	acatccgc agat	ttagga aacti	ttcata gagtg	taatc ttttc	tttct 720	0
gtaaaaatat at	tttacttg tagc	attgga gatti	tgttat gagat	tatct tactt	agcat 780	0
ttagtgaata at	ctattagc ctat	tttgcc gacg	tg		816	6